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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/619,047C

DATE: 05/11/2001

TIME: 13:55:55

Input Set : A:\CHEM1110.ST25.txt
 Output Set: N:\CRF3\05112001\I619047C.raw

P.S.

3 <110> APPLICANT: Chemicon International, Inc.
 4 LENG, Jay
 6 <120> TITLE OF INVENTION: PROTEASE SPECIFIC CLEAVABLE LUCIFERASES AND METHODS OF USE THEREOF
 8 <130> FILE REFERENCE: CHEM1110
 10 <140> CURRENT APPLICATION NUMBER: US 09/619,047C
 11 <141> CURRENT FILING DATE: 2000-07-18
 13 <160> NUMBER OF SEQ ID NOS: 29
 15 <170> SOFTWARE: PatentIn version 3.0
 17 <210> SEQ ID NO: 1
 18 <211> LENGTH: 936
 19 <212> TYPE: DNA
 20 <213> ORGANISM: Renilla reniformis
 22 <220> FEATURE:
 23 <221> NAME/KEY: CDS
 24 <222> LOCATION: (1)..(936)
 26 <400> SEQUENCE: 1
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 28 Met Thr Ser Lys Val Tyr Asp Pro Glu Gln Arg Lys Arg Met Ile Thr
 29 1 5 10 15
 31 ggt ccg cag tgg tgg gcc aga tgt aaa caa atg aat gtt ctt gat tca 96
 32 Gly Pro Gln Trp Trp Ala Arg Cys Lys Gln Met Asn Val Leu Asp Ser
 33 20 25 30
 35 ttt att aat tat tat gat tca gaa aaa cat gca gaa aat gct gtt att 144
 36 Phe Ile Asn Tyr Tyr Asp Ser Glu Lys His Ala Glu Asn Ala Val Ile
 37 35 40 45
 39 ttt tta cat ggt aac gcg gcc tct tct tat tta tgg cga cat gtt gtg 192
 40 Phe Leu His Gly Asn Ala Ala Ser Ser Tyr Leu Trp Arg His Val Val
 41 50 55 60
 43 cca cat att gag cca gta gcg cgg tgt att ata cca gat ctt att ggt 240
 44 Pro His Ile Glu Pro Val Ala Arg Cys Ile Ile Pro Asp Leu Ile Gly
 45 65 70 75 80
 47 atg ggc aaa tca ggc aaa tct ggt aat ggt tct tat agg tta ctt gat 288
 48 Met Gly Lys Ser Gly Lys Ser Gly Asn Gly Ser Tyr Arg Leu Leu Asp
 49 85 90 95
 51 cat tac aaa tat ctt act gca tgg ttt gaa ctt ctt aat tta cca aag 336
 52 His Tyr Lys Tyr Leu Thr Ala Trp Phe Glu Leu Leu Asn Leu Pro Lys
 53 100 105 110
 55 aag atc att ttt gtc ggc cat gat tgg ggt gct tgt ttg gca ttt cat 384
 56 Lys Ile Ile Phe Val Gly His Asp Trp Gly Ala Cys Leu Ala Phe His
 57 115 120 125
 59 tat agc tat gag cat caa gat aag atc aaa gca ata gtt cac gct gaa 432
 60 Tyr Ser Tyr Glu His Gln Asp Lys Ile Lys Ala Ile Val His Ala Glu
 61 130 135 140
 63 agt gta gta gat gtg att gaa tca tgg gat gaa tgg cct gat att gaa 480
 64 Ser Val Val Asp Val Ile Glu Ser Trp Asp Glu Trp Pro Asp Ile Glu
 65 145 150 155 160
 67 gaa gat att gcg ttg atc aaa tct gaa gaa gga gaa aaa atg gtt ttg 528

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68	Glu	Asp	Ile	Ala	Leu	Ile	Lys	Ser	Glu	Glu	Gly	Glu	Lys	Met	Val	Leu	
69				165					170					175			
71	gag	aat	aac	tcc	tcc	gtg	gaa	acc	atg	ttg	cca	tca	aaa	atc	atg	aga	576
72	Glu	Asn	Asn	Phe	Phe	Val	Glu	Thr	Met	Leu	Pro	Ser	Lys	Ile	Met	Arg	
73				180				185					190				
75	aag	tta	gaa	cca	gaa	gaa	ttt	gca	gca	tat	ctt	gaa	cca	ttc	aaa	gag	624
76	Lys	Leu	Glu	Pro	Glu	Glu	Phe	Ala	Ala	Tyr	Leu	Glu	Pro	Phe	Lys	Glu	
77				195				200				205					
79	aaa	ggg	gaa	gtt	cgt	cgt	cca	aca	tta	tca	tgg	cct	cgt	gaa	atc	ccg	672
80	Lys	Gly	Glu	Val	Arg	Arg	Pro	Thr	Leu	Ser	Trp	Pro	Arg	Glu	Ile	Pro	
81				210				215				220					
83	tta	gta	aaa	ggg	ggg	aaa	cct	gac	gtt	gta	caa	att	gtt	agg	aat	tat	720
84	Leu	Val	Lys	Gly	Gly	Lys	Pro	Asp	Val	Val	Gln	Ile	Val	Arg	Asn	Tyr	
85	225			230				235				235			240		
87	aat	gct	tat	cta	cgt	gca	agt	gat	gat	tta	cca	aaa	atg	ttt	att	gaa	768
88	Asn	Ala	Tyr	Leu	Arg	Ala	Ser	Asp	Asp	Leu	Pro	Lys	Met	Phe	Ile	Glu	
89				245				250				255					
91	tcg	gat	cca	gga	tcc	ttt	tcc	aat	gct	att	gtt	gaa	ggc	gcc	aag	aag	816
92	Ser	Asp	Pro	Gly	Phe	Phe	Ser	Asn	Ala	Ile	Val	Glu	Gly	Ala	Lys	Lys	
93				260				265				270					
95	ttt	cct	aat	act	gaa	ttt	gtc	aaa	gta	aaa	ggg	ctt	cat	ttt	tcg	caa	864
96	Phe	Pro	Asn	Thr	Glu	Phe	Val	Lys	Val	Lys	Gly	Leu	His	Phe	Ser	Gln	
97				275				280				285					
99	gaa	gat	gca	cct	gat	gaa	atg	gga	aaa	tat	atc	aaa	tcg	tcc	gtt	gag	912
100	Glu	Asp	Ala	Pro	Asp	Glu	Met	Gly	Lys	Tyr	Ile	Lys	Ser	Phe	Val	Glu	
101				290				295				300					936
103	cga	gtt	ctc	aaa	aat	gaa	caa	taa									
104	Arg	Val	Leu	Lys	Asn	Glu	Gln										
105	305				310												
108	<210>	SEQ ID NO:	2														
109	<211>	LENGTH:	311														
110	<212>	TYPE:	PRT														
111	<213>	ORGANISM:	Renilla reniformis														
113	<400>	SEQUENCE:	2														
115	Met	Thr	Ser	Lys	Val	Tyr	Asp	Pro	Glu	Gln	Arg	Lys	Arg	Met	Ile	Thr	
116	1				5				10					15			
119	Gly	Pro	Gln	Trp	Trp	Ala	Arg	Cys	Lys	Gln	Met	Asn	Val	Leu	Asp	Ser	
120					20				25				30				
123	Phe	Ile	Asn	Tyr	Tyr	Asp	Ser	Glu	Lys	His	Ala	Glu	Asn	Ala	Val	Ile	
124					35			40				45					
127	Phe	Leu	His	Gly	Asn	Ala	Ala	Ser	Ser	Tyr	Leu	Trp	Arg	His	Val	Val	
128					50			55			60						
131	Pro	His	Ile	Glu	Pro	Val	Ala	Arg	Cys	Ile	Ile	Pro	Asp	Leu	Ile	Gly	
132	65					70			75				80				
135	Met	Gly	Lys	Ser	Gly	Lys	Ser	Gly	Asn	Gly	Ser	Tyr	Arg	Leu	Leu	Asp	
136					85				90				95				
139	His	Tyr	Lys	Tyr	Leu	Thr	Ala	Trp	Phe	Glu	Leu	Leu	Asn	Leu	Pro	Lys	
140					100				105				110				
143	Lys	Ile	Ile	Phe	Val	Gly	His	Asp	Trp	Gly	Ala	Cys	Leu	Ala	Phe	His	

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144	115	120	125
147	Tyr Ser Tyr Glu His Gln Asp Lys Ile Lys Ala Ile Val His Ala Glu		
148	130	135	140
151	Ser Val Val Asp Val Ile Glu Ser Trp Asp Glu Trp Pro Asp Ile Glu		
152	145	150	155
155	Glu Asp Ile Ala Leu Ile Lys Ser Glu Glu Gly Glu Lys Met Val Leu		160
156	165	170	175
159	Glu Asn Asn Phe Phe Val Glu Thr Met Leu Pro Ser Lys Ile Met Arg		
160	180	185	190
163	Lys Leu Glu Pro Glu Glu Phe Ala Ala Tyr Leu Glu Pro Phe Lys Glu		
164	195	200	205
167	Lys Gly Glu Val Arg Arg Pro Thr Leu Ser Trp Pro Arg Glu Ile Pro		
168	210	215	220
171	Leu Val Lys Gly Gly Lys Pro Asp Val Val Gln Ile Val Arg Asn Tyr		
172	225	230	235
175	Asn Ala Tyr Leu Arg Ala Ser Asp Asp Leu Pro Lys Met Phe Ile Glu		240
176	245	250	255
179	Ser Asp Pro Gly Phe Phe Ser Asn Ala Ile Val Glu Gly Ala Lys Lys		
180	260	265	270
183	Phe Pro Asn Thr Glu Phe Val Lys Val Lys Gly Leu His Phe Ser Gln		
184	275	280	285
187	Glu Asp Ala Pro Asp Glu Met Gly Lys Tyr Ile Lys Ser Phe Val Glu		
188	290	295	300
191	Arg Val Leu Lys Asn Glu Gln		
192	305	310	
195	<210> SEQ ID NO: 3		
196	<211> LENGTH: 936		
197	<212> TYPE: DNA		
198	<213> ORGANISM: Renilla reniformis (mutated sequence)		
200	<220> FEATURE:		
201	<221> NAME/KEY: CDS		
202	<222> LOCATION: (1)..(936)		
204	<400> SEQUENCE: 3		
205	atg act tcg aaa gtt tat gat cca gaa caa agg aaa cgg atg ata act		48
206	Met Thr Ser Lys Val Tyr Asp Pro Glu Gln Arg Lys Arg Met Ile Thr		
207	1 5 10 15		
209	ggt ccg cag tgg tgg gcc aga tgt aaa caa atg aat gtt ctt gat tca		96
210	Gly Pro Gln Trp Trp Ala Arg Cys Lys Gln Met Asn Val Leu Asp Ser		
211	20 25 30		
213	ttt att aat tat tat gat tca gaa aaa cat gca gaa aat gct gtt att		144
214	Phe Ile Asn Tyr Tyr Asp Ser Glu Lys His Ala Glu Asn Ala Val Ile		
215	35 40 45		
217	ttt tta cat ggt aac gcg gcc tct tat tta tgg cga cat gtt gtg		192
218	Phe Leu His Gly Asn Ala Ala Ser Ser Tyr Leu Trp Arg His Val Val		
219	50 55 60		
221	cca cat att gag cca gta gcg cgg tgt att ata cca gat ctt att ggt		240
222	Pro His Ile Glu Pro Val Ala Arg Cys Ile Ile Pro Asp Leu Ile Gly		
223	65 70 75 80		
225	atg ggc aaa tca ggc aaa tct ggt aat ggt tct tat agg tta ctt gat		288

RAW SEQUENCE LISTING DATE: 05/11/2001
PATENT APPLICATION: US/09/619,047C TIME: 13:55:55

Input Set : A:\CHEM1110.ST25.txt
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PATENT APPLICATION: US/09/619,047C

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Input Set : A:\CHEM1110.ST25.txt
Output Set: N:\CRF3\05112001\I619047C.raw

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293 Met Thr Ser Lys Val Tyr Asp Pro Glu Gln Arg Lys Arg Met Ile Thr
294 1      5      10      15
297 Gly Pro Gln Trp Trp Ala Arg Cys Lys Gln Met Asn Val Leu Asp Ser
298      20      25      30
301 Phe Ile Asn Tyr Tyr Asp Ser Glu Lys His Ala Glu Asn Ala Val Ile
302      35      40      45
305 Phe Leu His Gly Asn Ala Ala Ser Ser Tyr Leu Trp Arg His Val Val
306      50      55      60
309 Pro His Ile Glu Pro Val Ala Arg Cys Ile Ile Pro Asp Leu Ile Gly
310 65      70      75      80
313 Met Gly Lys Ser Gly Lys Ser Gly Asn Gly Ser Tyr Arg Leu Leu Asp
314      85      90      95
317 His Tyr Lys Tyr Leu Thr Ala Trp Phe Glu Leu Leu Asn Leu Pro Lys
318      100     105     110
321 Lys Ile Ile Phe Val Gly His Asp Trp Gly Ala Cys Leu Ala Phe His
322      115     120     125
325 Tyr Ser Tyr Glu His Gln Asp Lys Ile Lys Ala Ile Val His Ala Glu
326      130     135     140
329 Ser Val Val Asp Val Ile Glu Ser Trp Asp Glu Trp Pro Asp Ile Glu
330 145     150     155     160
333 Glu Asp Ile Ala Leu Ile Lys Ser Glu Glu Gly Glu Lys Met Val Leu
334      165     170     175
337 Glu Asn Asn Phe Phe Val Glu Thr Met Leu Pro Ser Lys Ile Met Arg
338      180     185     190
341 Lys Leu Glu Pro Asp Glu Val Asp Ala Tyr Leu Glu Pro Phe Lys Glu
342      195     200     205
345 Lys Gly Glu Val Arg Arg Pro Thr Leu Ser Trp Pro Arg Glu Ile Pro
346      210     215     220
349 Leu Val Lys Gly Gly Lys Pro Asp Val Val Gln Ile Val Arg Asn Tyr
350 225     230     235     240
353 Asn Ala Tyr Leu Arg Ala Ser Asp Asp Leu Pro Lys Met Phe Ile Glu
354      245     250     255
357 Ser Asp Pro Gly Phe Phe Ser Asn Ala Ile Val Glu Gly Ala Lys Lys
358      260     265     270
361 Phe Pro Asn Thr Glu Phe Val Lys Val Lys Gly Leu His Phe Ser Gln
362      275     280     285
365 Glu Asp Ala Pro Asp Glu Met Gly Lys Tyr Ile Lys Ser Phe Val Glu
366      290     295     300
369 Arg Val Leu Lys Asn Glu Gln
370 305     310
373 <210> SEQ ID NO: 5
374 <211> LENGTH: 8
375 <212> TYPE: PRT
376 <213> ORGANISM: Artificial sequence
378 <220> FEATURE:
379 <223> OTHER INFORMATION: Description of Artificial sequence: Protease recognition sequence
381 <400> SEQUENCE: 5
383 Ser Gln Asn Tyr Pro Ile Val Gln
384 1      5

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Please Note:

Use f n and/ r Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/619,047C

DATE: 05/11/2001
TIME: 13:55:56

Input Set : A:\CHEM1110.ST25.txt
Output Set: N:\CRF3\05112001\I619047C.raw

L:613 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:632 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:677 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26